

## EAST Search History

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	0	pyruvate adj decarboxylase and ethanol and yeast and gene and bacillus	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:23
L2	0	pyruvate adj decarboxylase and ethanol	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:23
L3	3478	pyruvate adj decarboxylase and ethanol	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:23
L4	1073	I3 and gene and bacillus	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:25
L5	924	I3 and gene and bacillus and alcohol adj dehydrogenase	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:26
L6	431	I5 and gram-positive	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:27
L7	29	I5 and gram-positive and lactate adj dehydrogenase	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:27

and display fields  
 NEWS 12 JUN 28 Price changes in full-text patent databases EPFULL and PCTFULL  
 NEWS 13 JUL 11 CHEMSAFE reloaded and enhanced  
 NEWS 14 JUL 14 FSTA enhanced with Japanese patents  
 NEWS 15 JUL 19 Coverage of Research Disclosure reinstated in DWPI

NEWS EXPRESS JUNE 30 CURRENT WINDOWS VERSION IS V8.01b, CURRENT  
 MACINTOSH VERSION IS V6.0c(ENG) AND V6.0Jc(JP),  
 AND CURRENT DISCOVER FILE IS DATED 26 JUNE 2006.

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=> s ethanol and yeast and gene  
 L1 3636 ETHANOL AND YEAST AND GENE

=> s l1 and bacillus  
 L2 90 L1 AND BACILLUS

=> s l2 and pyruvate (w) decarboxylase  
L3 7 L2 AND PYRUVATE (W) DECARBOXYLASE

=> d ibib abs l3 1-7

L3 ANSWER 1 OF 7 MEDLINE on STN  
ACCESSION NUMBER: 2005658391 MEDLINE  
DOCUMENT NUMBER: PubMed ID: 16339947  
TITLE: Construction and expression of an ethanol  
production operon in Gram-positive bacteria.  
AUTHOR: Talarico Lee A; Gil Malgorzata A; Yomano Lorraine P; Ingram  
Lonnie O; Maupin-Furlow Julie A  
CORPORATE SOURCE: Department of Microbiology and Cell Science, University of  
Florida, Gainesville, FL 32611-0700, USA.  
SOURCE: Microbiology (Reading, England), (2005 Dec) Vol. 151, No.  
Pt 12, pp. 4023-31.  
Journal code: 9430468. ISSN: 1350-0872.  
PUB. COUNTRY: England; United Kingdom  
DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)  
LANGUAGE: English  
FILE SEGMENT: Priority Journals  
ENTRY MONTH: 200602  
ENTRY DATE: Entered STN: 18 Dec 2005  
Last Updated on STN: 1 Mar 2006  
Entered Medline: 28 Feb 2006

AB Pyruvate decarboxylase (PDC), an enzyme central to  
homoethanol fermentation, catalyses the non-oxidative decarboxylation of  
pyruvate to acetaldehyde with release of carbon dioxide. PDC enzymes from  
diverse organisms have different kinetic properties, thermal stability and  
codon usage that are likely to offer unique advantages for the development  
of desirable Gram-positive biocatalysts for use in the ethanol  
industry. To examine this further, pdc genes from bacteria to  
yeast were expressed in the Gram-positive host *Bacillus*  
*megaterium*. The PDC activity and protein levels were determined for each  
strain. In addition, the levels of pdc-specific mRNA transcripts and  
stability of recombinant proteins were assessed. From this analysis, the  
pdc gene of Gram-positive *Sarcina ventriculi* was found to be the  
most advantageous for engineering high-level synthesis of PDC in a  
Gram-positive host. This gene was thus selected for  
transcriptional coupling to the alcohol dehydrogenase gene (*adh*)  
of *Geobacillus stearothermophilus*. The resulting Gram-positive  
ethanol production operon was expressed at high levels in *B.*  
*megaterium*. Extracts from this recombinant were shown to catalyse the  
production of ethanol from pyruvate.

L3 ANSWER 2 OF 7 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN  
ACCESSION NUMBER: 2006:151966 BIOSIS  
DOCUMENT NUMBER: PREV200600152190  
TITLE: Construction and expression of an ethanol  
production operon in Gram-positive bacteria.  
AUTHOR(S): Talarico, Lee A.; Gil, Malgorzata A.; Yomano, Lorraine P.;  
Ingram, Lonnie O.; Maupin-Furlow, Julie A. [Reprint Author]  
CORPORATE SOURCE: Univ Florida, Dept Microbiol and Cell Sci, Gainesville, FL  
32611 USA  
jmaupin@ufl.edu  
SOURCE: Microbiology (Reading), (DEC 2005) Vol. 151, No. Part 12,  
pp. 4023-4031.  
ISSN: 1350-0872.  
DOCUMENT TYPE: Article  
LANGUAGE: English  
ENTRY DATE: Entered STN: 1 Mar 2006  
Last Updated on STN: 1 Mar 2006

AB Pyruvate decarboxylase (PDC), an enzyme central to  
homoethanol fermentation, catalyses the non-oxidative decarboxylation of

pyruvate to acetaldehyde with release of carbon dioxide. 1 enzymes from diverse organisms have different kinetic properties, thermal stability and codon usage that are likely to offer unique advantages for the development of desirable Gram-positive biocatalysts for use in the ethanol industry. To examine this further, pdc genes from bacteria to yeast were expressed in the Gram-positive host *Bacillus megaterium*. The PDC activity and protein levels were determined for each strain. In addition, the levels of pdc-specific mRNA transcripts and stability of recombinant proteins were assessed. From this analysis, the pdc gene of Gram-positive *Sarcina ventriculi* was found to be the most advantageous for engineering high-level synthesis of PDC in a Gram-positive host. This gene was thus selected for transcriptional coupling to the alcohol dehydrogenase gene (adh) of *Geobacillus stearothermophilus*. The resulting Gram-positive ethanol production operon was expressed at high levels in *B. megaterium*. Extracts from this recombinant were shown to catalyse the production of ethanol from pyruvate.

L3 ANSWER 3 OF 7 CAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER: 2006:17838 CAPLUS  
DOCUMENT NUMBER: 144:405166  
TITLE: Construction and expression of an ethanol production operon in Gram-positive bacteria  
AUTHOR(S): Talarico, Lee A.; Gil, Malgorzata A.; Yomano, Lorraine P.; Ingram, Lonnie O.; Maupin-Furlow, Julie A.  
CORPORATE SOURCE: Department of Microbiology and Cell Science, University of Florida, Gainesville, FL, 32611-0700, USA  
SOURCE: Microbiology (Reading, United Kingdom) (2005), 151(12), 4023-4031  
CODEN: MROBEO; ISSN: 1350-0872  
PUBLISHER: Society for General Microbiology  
DOCUMENT TYPE: Journal  
LANGUAGE: English

AB Pyruvate decarboxylase (PDC), an enzyme central to homoethanol fermentation, catalyzes the non-oxidative decarboxylation of pyruvate to acetaldehyde with release of carbon dioxide. PDC enzymes from diverse organisms have different kinetic properties, thermal stability and codon usage that are likely to offer unique advantages for the development of desirable Gram-pos. biocatalysts for use in the ethanol industry. To examine this further, pdc genes from bacteria to yeast were expressed in the Gram-pos. host *Bacillus megaterium*. The PDC activity and protein levels were determined for each strain. In addition, the levels of pdc-specific mRNA transcripts and stability of recombinant proteins were assessed. From this anal., the pdc gene of Gram-pos. *Sarcina ventriculi* was found to be the most advantageous for engineering high-level synthesis of PDC in a Gram-pos. host. This gene was thus selected for transcriptional coupling to the alc. dehydrogenase gene (adh) of *Geobacillus stearothermophilus*. The resulting Gram-pos. ethanol production operon was expressed at high levels in *B. megaterium*. Exts. from this recombinant were shown to catalyze the production of ethanol from pyruvate.

REFERENCE COUNT: 35 THERE ARE 35 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L3 ANSWER 4 OF 7 CAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER: 1992:649986 CAPLUS  
DOCUMENT NUMBER: 117:249986  
TITLE: Ethanol production by by bacteria carrying foreign genes for alcohol dehydrogenase and pyruvate decarboxylase  
INVENTOR(S): Ingram, Lonnie O.; Beall, David S.; Burchhardt, Gerhard F. H.; Guimaraes, Walter V.; Ohta, Kazuyoshi;

PATENT ASSIGNEE(S): Wood, Brent E.; Shanmugam, Keelnatham T.; Fowler, David A.; Ben-Bassat, Arie  
 SOURCE: PCT Int. Appl., 153 pp.  
 CODEN: PIXXD2  
 DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 10  
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9216615	A1	19921001	WO 1992-US1807	19920318
W: AT, AU, BB, BG, BR, CA, CH, CS, DE, DK, ES, FI, GB, HU, JP, KP, KR, LK, LU, MG, MN, MW, NL, NO, PL, RO, RU, SD, SE, US				
RW: AT, BE, BF, BJ, CF, CG, CH, CI, CM, DE, DK, ES, FR, GA, GB, GN, GR, IT, LU, MC, ML, MR, NL, SE, SN, TD, TG				
US 5424202	A	19950613	US 1992-846344	19920306
AU 9217794	A1	19921021	AU 1992-17794	19920318
AU 672748	B2	19961017		
CN 1070424	A	19930331	CN 1992-101877	19920318
CN 1065915	B	20010516		
EP 576621	A1	19940105	EP 1992-910933	19920318
EP 576621	B1	20010228		
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, MC, NL, SE				
JP 06505875	T2	19940707	JP 1992-509941	19920318
JP 3457664	B2	20031020		
BR 9205782	A	19940726	BR 1992-5782	19920318
AT 199389	E	20010315	AT 1992-910933	19920318
NO 9303178	A	19931108	NO 1993-3178	19930907
NO 315567	B1	20030922		
CN 1342773	A	20020403	CN 2000-131779	20001020
AU 2005248924	A1	20060202	AU 2005-248924	20051223

PRIORITY APPLN. INFO.:  
 US 1991-670821 A 19910318  
 US 1992-846344 A 19920306  
 US 1988-239099 B2 19880831  
 US 1989-352062 A2 19890515  
 US 1990-624277 B2 19901207  
 WO 1992-US1807 A 19920318  
 AU 2002-35569 A3 20020419

AB Bacterial hosts, excluding *Escherichia coli*, expressing heterologous genes for alc. dehydrogenase (I) and pyruvate decarboxylase (II) are used for manufacture of EtOH. II is used to prevent accumulation of acid metabolites. Plasmids, e.g. pLOI555 carrying genes for I and II of *Zymomonas mobilis* driven by the lac promoter, are provided for preparation of the host. The method is further improved by transforming the host with genes for proteins that facilitate transport and metabolism of oligosaccharides, e.g., of C5-6 sugars, which host is, preferably, also expressing a heterologous gene for a polysaccharase such as a cellulolytic enzyme, a xylanolytic enzyme, or a starch-degrading enzyme. These hosts also preferably express heterologous genes for polysaccharide-degrading enzymes (e.g. those degrading cellulose, xylans, or starch). A cost-effective fermentation process for manufacturing EtOH from oligosaccharide feedstocks using a single, genetically engineered microorganism is also disclosed. An ethanologenic strain *Klebsiella oxytoca* M5A1(pLOI555) was prepared and was further transformed with plasmid pLOI2003 encoding xylanase (gene xynZ) and xylosidase (gene xylB) of *Clostridium thermocellum* to obtain a transformant capable of converting xylan to EtOH.

L3 ANSWER 5 OF 7 LIFESCI COPYRIGHT 2006 CSA on STN  
 ACCESSION NUMBER: 2006:26910 LIFESCI  
 TITLE: Construction and expression of an ethanol

production operon in Gram-positive bacteria  
 AUTHOR: Talarico, Lee A.; Gil, Malgorzata A.; Yomano, Lorraine P.;  
 Ingram, Lonnie O.; Maupin-Furrow, Julie A.  
 CORPORATE SOURCE: Department of Microbiology and Cell Science, University of  
 Florida, Gainesville, FL 32611-0700, USA; E-mail:  
 jmaupin@ufl.edu  
 SOURCE: Microbiology, (20051200) vol. 151, no. 12, pp. 4023-4031.  
 ISSN: 1350-0872.  
 DOCUMENT TYPE: Journal  
 FILE SEGMENT: W2; J  
 LANGUAGE: English  
 SUMMARY LANGUAGE: English

AB Pyruvate decarboxylase (PDC), an enzyme central to  
 homoethanol fermentation, catalyses the non-oxidative decarboxylation of  
 pyruvate to acetaldehyde with release of carbon dioxide. PDC enzymes from  
 diverse organisms have different kinetic properties, thermal stability and  
 codon usage that are likely to offer unique advantages for the development  
 of desirable Gram-positive biocatalysts for use in the ethanol  
 industry. To examine this further, pdc genes from bacteria to  
 yeast were expressed in the Gram-positive host *Bacillus*  
*megaterium*. The PDC activity and protein levels were determined for each  
 strain. In addition, the levels of pdc-specific mRNA transcripts and  
 stability of recombinant proteins were assessed. From this analysis, the  
 pdc gene of Gram-positive *Sarcina ventriculi* was found to be the  
 most advantageous for engineering high-level synthesis of PDC in a  
 Gram-positive host. This gene was thus selected for  
 transcriptional coupling to the alcohol dehydrogenase gene (adh)  
 of *Geobacillus stearothermophilus*. The resulting Gram-positive  
 ethanol production operon was expressed at high levels in *B.*  
*megaterium*. Extracts from this recombinant were shown to catalyse the  
 production of ethanol from pyruvate.

L3 ANSWER 6 OF 7 LIFESCI COPYRIGHT 2006 CSA on STN  
 ACCESSION NUMBER: 84:75038 LIFESCI  
 TITLE: Xylose-isomerase in yeast.  
 AUTHOR: Wilhelm, M.; Erhart, E.; Hollenberg, C.P.  
 CORPORATE SOURCE: Inst. Mikrobiol., Univ. Duesseldorf, Duesseldorf, FRG  
 SOURCE: RIV. BIOL., (1984) vol. 77, no. 4, pp. 607-608.  
 Meeting Info.: International Course on Microbial Breeding.  
 Spoleto (Italy). 3-8 Sep 1984.  
 DOCUMENT TYPE: Journal  
 TREATMENT CODE: Conference  
 FILE SEGMENT: G; W; N; K  
 LANGUAGE: English

AB *Saccharomyces cerevisiae* is not able to ferment xylose to ethanol  
 . To explore the feasibility of constructing yeast strains that  
 can produce ethanol from xylose, the authors have decided to  
 introduce a xylose isomerase gene into *S. cerevisiae* under  
 control of the yeast pyruvate decarboxylase  
 promoter and to study the effects of its expression. As a first step the  
 authors have isolated a BamHI fragment from *Bacillus subtilis*  
 which carries the genetic information for xylose isomerase and  
 exlyulokinase.

L3 ANSWER 7 OF 7 EMBASE COPYRIGHT (c) 2006 Elsevier B.V. All rights  
 reserved on STN  
 ACCESSION NUMBER: 2005583341 EMBASE  
 TITLE: Construction and expression of an ethanol  
 production operon in Gram-positive bacteria.  
 AUTHOR: Talarico L.A.; Gil M.A.; Yomano L.P.; Ingram L.O.;  
 Maupin-Furrow J.A.  
 CORPORATE SOURCE: J.A. Maupin-Furrow, Department of Microbiology and Cell  
 Science, University of Florida, Gainesville, FL 32611-0700,  
 United States. jmaupin@ufl.edu

SOURCE: Microbiology, (2005) Vol. 151, No. 12, pp. 4023-4031. .  
 Refs: 35  
 ISSN: 1350-0872 CODEN: MROBEO  
 COUNTRY: United Kingdom  
 DOCUMENT TYPE: Journal; Article  
 FILE SEGMENT: 004 Microbiology  
 LANGUAGE: English  
 SUMMARY LANGUAGE: English  
 ENTRY DATE: Entered STN: 12 Jan 2006  
 Last Updated on STN: 12 Jan 2006

AB Pyruvate decarboxylase (PDC), an enzyme central to homoethanol fermentation, catalyses the non-oxidative decarboxylation of pyruvate to acetaldehyde with release of carbon dioxide. PDC enzymes from diverse organisms have different kinetic properties, thermal stability and codon usage that are likely to offer unique advantages for the development of desirable Gram-positive biocatalysts for use in the ethanol industry. To examine this further, pdc genes from bacteria to yeast were expressed in the Gram-positive host *Bacillus megaterium*. The PDC activity and protein levels were determined for each strain. In addition, the levels of pdc-specific mRNA transcripts and stability of recombinant proteins were assessed. From this analysis, the pdc gene of Gram-positive *Sarcina ventriculi* was found to be the most advantageous for engineering high-level synthesis of PDC in a Gram-positive host. This gene was thus selected for transcriptional coupling to the alcohol dehydrogenase gene (adh) of *Geobacillus stearothermophilus*. The resulting Gram-positive ethanol production operon was expressed at high levels in *B. megaterium*. Extracts from this recombinant were shown to catalyse the production of ethanol from pyruvate. .COPYRG. 2005 SGM.

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